

<u>SEQUENCE LISTING</u> (37 C.F.R. §§ 1.821 - 1.825)

- (1) GENERAL INFORMATION:
- (i) APPLICANT: ROBERT WEBBER
- (ii) TITLE OF INVENTION: IMMUNOASSAY METHOD EMPLOYING MONOCLONAL ANTIBODY REACTIVE TO HUMAN INOS
- (iii) NUMBER OF SEQUENCES: 126
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: BIELEN, PETERSON & LAMPE
 - (B) STREET: 1990 N. CALIFORNIA BOULEVARD, SUITE 720
 - (C) CITY: WALNUT CREEK
 - (D) STATE: CALIFORNIA
 - (E) COUNTRY: UNITED STATES OF AMERICA
 - (F) ZIP: 94596
- (V) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: DISKETTE 3.5 INCH, 1.44 MB FOR FORMATTED
 - (B) COMPUTER: IBM PC COMPATIBLE
 - (C) OPERATING SYSTEM: DOS
 - (D) SOFTWARE: WORDPERFECT 5.1
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: NONE
 - (B) FILING DATE: NONE
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/634,332
 - (B) FILING DATE: 12 APRIL 1996
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: THEODORE J. BIELEN, JR.
 - (B) REGISTRATION NUMBER: 27,420
 - (C) REFERENCE/DOCKET NUMBER: 12280
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (925) 937-1515
 - (B) TELEFAX: (925) 937-1529
 - (2) INFORMATION FOR SEQ ID NO: 1:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (25-42)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
- Asn Asn Asn Val Glu Lys Ala Pro Cys Ala Thr Ser Ser 5

Pro Val Thr Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 2:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: MOUSE INOS (25-42)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Asn Asn Asn Val Lys Lys Thr Pro Cys Ala Val Leu Ser 5 10

Pro Thr Ile Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 3:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: RAT iNOS (25-42)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Asn Asn Asn Val Glu Lys Thr Pro Gly Ala Ile Pro Ser

Pro Thr Thr Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 4:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE

- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-54)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Pro Val Thr Gln Asp Asp Leu Gln Tyr His Ser 5 10 Gln Gln Asn Ser Lys 15

- (2) INFORMATION FOR SEQ ID NO: 5:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (781-798)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Ala Leu Val Gln Gly Ile Leu Glu Arg Val Pro Val 5 10 Gly His Pro Thr Pro

- Gly Pro Thr Pro His
 - (2) INFORMATION FOR SEQ ID NO: 6:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 17
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: MOUSE INOS (776-792)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Xaa Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10 Cys Pro Thr Pro His

15

(2) INFORMATION FOR SEQ ID NO: 7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: RAT iNOS (780-794) (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7: Gln Gly Ile Leu Glu Arg Val Val Xaa Xaa Leu Val Asp 10 5 Cys Ser Ser Pro Xaa 15 (2) INFORMATION FOR SEQ ID NO: 8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN INOS (985-1002) (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8: Phe Arg Ser Phe Gly Ile Val Pro Trp Gln Gln Arg Leu 10 His Gln Ser His Asp 15 (2) INFORMATION FOR SEQ ID NO: 9: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) **FEATURE:** (A) NAME/KEY: MOUSE iNOS (978-995)

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

SEQUENCE DESCRIPTION: SEQ ID NO: 9:

(B) LOCATION:

(xi)

(D) OTHER INFORMATION:

- Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
 5 10
 His Asp Ser Gln His
 15
 - (2) INFORMATION FOR SEQ ID NO: 10:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: RAT iNOS (982-998)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
- Ala Phe Trp Gly Ile Pro Arg Ser Phe Gln Gln Arg Leu 5 10 His Ser Gln His Asp 15
 - (2) INFORMATION FOR SEQ ID NO: 11:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN nNOS (1256-1273)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
- Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Gln
 5 10
 Phe Asp Ile Gln His
 15
 - (2) INFORMATION FOR SEQ ID NO: 12:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN eNOS (1017-1031)

- (B) LOCATION:
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu
 5 10
- His Asp Xaa Xaa Xaa 15
 - (2) INFORMATION FOR SEQ ID NO: 13:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: BOVINE eNOS (1019-1033)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu 5 10
- His Asp Xaa Xaa Xaa 15
 - (2) INFORMATION FOR SEQ ID NO: 14:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (1009-1026)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
- Arg Met Thr Leu Val Phe Gly Cys Arg Arg Pro Asp Glu 5
- Asp His Ile Tyr Gln 15
 - (2) INFORMATION FOR SEQ ID NO: 15:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18

- (B) TYPE: AMINO ACID
- (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: RAT iNOS (1006-1023)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

Arg Met Thr Leu Val Phe Gly Cys Arg His Pro Glu Glu 5

Asp His Leu Tyr Gln 15

- (2) INFORMATION FOR SEQ ID NO: 16:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: MOUSE INOS (1002-1019)
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

Arg Met Ser Leu Val Phe Gly Cys Arg His Pro Glu Glu
5

Asp His Leu Tyr Gln 15

- (2) INFORMATION FOR SEQ ID NO: 17:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hnNOS $[2-16, Cys^{17}]$
 - (B) LOCATION: HUMAN NNOS: AMINO TERMINAL
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

Glu Asp His Met Phe Gly Val Gln Gln Ile Gln Pro Asn 5 10

Val Ile Cys 15

- (2) INFORMATION FOR SEQ ID NO: 18:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hnNOS [Cys¹⁴¹⁰-1411-1433]
 - (B) LOCATION: HUMAN NNOS: CARBOXYL TERMINAL
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
- Cys Arg Leu Arg Ser Glu Ser Ile Ala Phe Ile Glu Glu
 5 10
- Ser Lys Lys Asp Thr Asp Glu Val Phe Ser Ser 15 20
 - (2) INFORMATION FOR SEQ ID NO: 19:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hinos $[2-21, Ser^2]$
 - (B) LOCATION: HUMAN INOS: AMINO TERMINAL
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
- Ala Ser Pro Trp Lys Phe Leu Phe Lys Thr Lys Phe His
 5
- Gln Tyr Ala Met Asn Gly Glu 15 20
 - (2) INFORMATION FOR SEQ ID NO: 20:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hinos $[Cys^{1136}-1137-1153]$
 - (B) LOCATION: HUMAN INOS: CARBOXYL TERMINAL
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

Cys Lys Lys Asp Arg Val Ala Val Gln Pro Ser Ser Leu 5 10 Glu Met Ser Ala Leu 15

- (2) INFORMATION FOR SEQ ID NO: 21:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: heNOS [Cap-2-12, Cys¹³]
 - (B) LOCATION: HUMAN ENOS: AMINO TERMINAL WITH CAPROIC ACID ATTACHED
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
- Gly Asn Leu Lys Ser Val Ala Gln Glu Pro Gly Cys
 5 10
 - (2) INFORMATION FOR SEQ ID NO: 22:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: henos $[2-12, Cys^{13}]$
 - (B) LOCATION: HUMAN ENOS: AMINO TERMINAL WITHOUT CAPROIC ACID ATTACHED
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
- Gly Asn Leu Lys Ser Val Ala Gln Glu Pro Gly Cys
 5
 - (2) INFORMATION FOR SEQ ID NO: 23:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: heNOS [Cys 1181 -1182-1203]
 - (B) LOCATION: HUMAN eNOS: CARBOXYL TERMINAL
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

Cys Glu Arg Gln Leu Arg Glu Ala Val Pro Trp Ala Phe
5

Asp Pro Pro Gly Ser Asp Thr Asn Ser Pro 15 20

- (2) INFORMATION FOR SEQ ID NO: 24:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hiNOS [985-1002]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
 5 10
 His Asp Ser Gln His
- His Asp Ser Gln His
 - (2) INFORMATION FOR SEQ ID NO: 25:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hinos [985-1002]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
 5 10
 His Asp Ser Gln His
- His Asp Ser Gln His
 - (2) INFORMATION FOR SEQ ID NO: 26:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR

- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hiNOS [37-54]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:
- Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu
 5 10
- Ser Lys Gln Gln Asn 15
 - (2) INFORMATION FOR SEQ ID NO: 27:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hiNOS [781-798]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:
- Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp
- Gly Pro Thr Pro His
 - (2) INFORMATION FOR SEQ ID NO: 28:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hinos [25-42]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:
- Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser Ser
- Pro Val Thr Gln Asp

15

- (2) INFORMATION FOR SEQ ID NO: 29:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hinos [37-54]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:
- Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu
 5 10
 Ser Lys Gln Gln Asp
- Ser Lys Gln Gln Asn 15
 - (2) INFORMATION FOR SEQ ID NO: 30:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hiNOS [781-798]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:
- Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp
 5 10
- Gly Pro Thr Pro His
 - (2) INFORMATION FOR SEQ ID NO: 31:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: hiNOS [1009-1026]
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

Arq Met Thr Leu Val Phe Gly Ser Arg Arg Pro Asp Glu 5 10 His Tyr Gln Asp Ile 15

- (2) INFORMATION FOR SEQ ID NO: 32:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: (A3) LOCUS HUMAN INOS (25-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser Ser 10 Thr Pro Val Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 33:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: MOUSE INOS (25-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

Asn Val Lys Lys Thr Pro Ser Ala Val Asn Asn Leu 5 10 Pro Thr Ile Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 34:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: RAT iNOS (25-42)

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

Asn Asn Asn Val Glu Lys Thr Pro Gly Ala Ile Pro Ser 5

Pro Thr Thr Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 35:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (28-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

Val Glu Lys Ala Pro Ser Ala Thr Ser Ser Pro Val Thr
5 10

Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 36:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (31-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

Ala Pro Ser Ala Thr Ser Ser Pro Val Thr Gln Asp 5 10

- (2) INFORMATION FOR SEQ ID NO: 37:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR

MOLECULE TYPE: PEPTIDE (ii)

(ix) FEATURE:

- (A) NAME/KEY: HUMAN iNOS (34-42)
- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

Ala Ser Pro Val Thr Gln Asp Thr Ser 5

- (2) INFORMATION FOR SEQ ID NO: 38:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- MOLECULE TYPE: PEPTIDE (ii)
- (ix) **FEATURE:**
 - (A) NAME/KEY: HUMAN iNOS (37-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- SEQUENCE DESCRIPTION: SEQ ID NO: 38: (xi)

Pro Val Thr Gln Asp 5

- (2) INFORMATION FOR SEQ ID NO: 39:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (25-39)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser Ser 5 10

Pro Val

15

- (2) INFORMATION FOR SEQ ID NO: 40:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID

- (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (25-36)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40:
- Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser 5 10
 - (2) INFORMATION FOR SEQ ID NO: 41:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:

(ii)

(ix)

- (A) NAME/KEY: HUMAN iNOS (25-33)
- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41:
- Asn Asn Asn Val Glu Lys Ala Pro Ser 5
 - (2) INFORMATION FOR SEQ ID NO: 42:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (25-30)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

Asn Asn Asn Val Glu Lys 5

- (2) INFORMATION FOR SEQ ID NO: 43:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR

(ii) MOLECULE TYPE: PEPTIDE

(ix) FEATURE:

- (A) NAME/KEY: (A4) LOCUS HUMAN INOS (37-54)
- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43:

Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu
5
10

Ser Lys Gln Gln Asn 15

- (2) INFORMATION FOR SEQ ID NO: 44:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (40-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44:

Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Lys Gln 5 10

Gln Asn 15

- (2) INFORMATION FOR SEQ ID NO: 45:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (43-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 45:

Asp Leu Gln Tyr His Asn Leu Ser Lys Gln Gln Asn 5 10

(2) INFORMATION FOR SEQ ID NO: 46: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (46-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 46: Leu Ser Lys Gln Gln Asn Tyr His Asn 5 (2) INFORMATION FOR SEQ ID NO: 47: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN INOS (49-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 47: Leu Ser Lys Gln Gln Asn 5 (2) INFORMATION FOR SEQ ID NO: 48: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-51) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 48: Gln His Asn Ser Pro Val Thr Asp Asp Leu Gln Tyr 5

Ser

Lys 15

10

- (2) INFORMATION FOR SEQ ID NO: 49:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-48)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 49:
- Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn 5 10
 - (2) INFORMATION FOR SEQ ID NO: 50:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-45)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 50:
- Ser Pro Val Thr Gln Asp Asp Leu Gln 5
 - (2) INFORMATION FOR SEQ ID NO: 51:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 51:

Ser Pro Val Thr Gln Asp

- (2) INFORMATION FOR SEQ ID NO: 52:
 (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: (F6) LOCUS HUMAN INOS (781-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 52:
- Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10
- Gly Pro Thr Pro His
 - (2) INFORMATION FOR SEQ ID NO: 53:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN eNOS (806-824)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 53:
- Pro Gly Leu Val Glu Ala Leu Leu Ser Arg Val Glu Asp
 5 10
 Pro Pro Ala Pro Thr Clu
- Pro Pro Ala Pro Thr Glu 15
 - (2) INFORMATION FOR SEQ ID NO: 54:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (784-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 54:

Val Gln Gly Ile Leu Glu Arg Val Val Asp Gly Pro Thr
5 10
Pro His
15

- (2) INFORMATION FOR SEQ ID NO: 55:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (787-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 55:
- Ile Leu Glu Arg Val Val Asp Gly Pro Thr Pro His
 5 10
 - (2) INFORMATION FOR SEQ ID NO: 56:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (790-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 56:
- Arg Val Val Asp Gly Pro Thr Pro His
 5
 - (2) INFORMATION FOR SEQ ID NO: 57:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (793-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 57:

Asp Gly Pro Thr Pro His 5

- (2) INFORMATION FOR SEQ ID NO: 58:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (781-794)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 58:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10 Gly

- (2) INFORMATION FOR SEQ ID NO: 59:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (781-792)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 59:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val 5

- (2) INFORMATION FOR SEQ ID NO: 60:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (781-789)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 60:

Pro Ala Leu Val Gln Gly Ile Leu Glu 5

- (2) INFORMATION FOR SEQ ID NO: 61:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (781-786)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 61:

Pro Ala Leu Val Gln Gly 5

- (2) INFORMATION FOR SEQ ID NO: 62:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: (G11) LOCUS HUMAN INOS (985-1002)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 62:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
 5 10
- His Asp Ser Gln His
 - (2) INFORMATION FOR SEQ ID NO: 63:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN nNOS (1256-1273)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 63:

Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Gln
5 10
Phe Asp Ile Gln His
15

- (2) INFORMATION FOR SEQ ID NO: 64:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN eNOS (1017-1031)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 64:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu 5 10
- His Asp 15
 - (2) INFORMATION FOR SEQ ID NO: 65:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (988-1002)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 65:
- Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu His Asp Ser 5
- Gln His
 - (2) INFORMATION FOR SEQ ID NO: 66:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (991-1002)

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 66:
- Ser Phe Trp Gln Gln Arg Leu His Asp Ser Gln His 5
 - (2) INFORMATION FOR SEQ ID NO: 67:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (994-1002)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 67:
- Gln Gln Arg Leu His Asp Ser Gln His
 5
 - (2) INFORMATION FOR SEQ ID NO: 68:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (997-1002)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 68:
- His Asp Ser Gln His 5
 - (2) INFORMATION FOR SEQ ID NO: 69:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (985-998)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 69: Phe Arg Ser Phe Trp Gln Gln Arg Leu Gly Ile Val Pro 5 10 His Asp 15 (2) INFORMATION FOR SEQ ID NO: 70: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN iNOS (985-996) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 70: Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg 5 (2) INFORMATION FOR SEQ ID NO: 71: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-993) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 71: (xi) Gly Ile Val Phe Arg Ser Phe Trp Pro 5 (2) INFORMATION FOR SEQ ID NO: 72: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-990)

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 72:
- Gly Ile Val Pro Phe Arg
 5
 - (2) INFORMATION FOR SEQ ID NO: 73:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: (H1) LOCUS HUMAN INOS (1009-1026)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 73:

Arg Met Thr Leu Val Phe Gly Ser Arg Arg Pro Asp Glu
5
10

Asp His Ile Tyr Gln 15

- (2) INFORMATION FOR SEQ ID NO: 74:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 17
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN eNOS (1041-1057)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 74:

Met Thr Leu Val Phe Gly Ser Arg Ser Ser Gln Leu Asp

His Leu Tyr Arg

- (2) INFORMATION FOR SEQ ID NO: 75:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 17
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR

- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN nNOS (1281-1297)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 75:
- Met Val Leu Val Phe Gly Ser Arg Gln Ser Lys Ile Asp 5
- His Ile Tyr Arg
 - (2) INFORMATION FOR SEQ ID NO: 76:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (1012-1026)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 76:
- Leu Val Phe Gly Ser Arg Arg Pro Asp Glu Asp His Ile 5
- Tyr Gln 15
 - (2) INFORMATION FOR SEQ ID NO: 77:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (1015-1026)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 77:
- Gly Ser Arg Arg Pro Asp Glu Asp His Ile Tyr Glr 5

(2) INFORMATION FOR SEQ ID NO: 78: (i)SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1018-1026) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 78: Arq Pro Asp Glu Asp His Ile Tyr Gln 5 (2) INFORMATION FOR SEQ ID NO: 79: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN INOS (1021-1026) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 79: (xi) Glu Asp His Ile Tyr Gln (2) INFORMATION FOR SEQ ID NO: 80: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1009-1023) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQ ID NO: 80: SEQUENCE DESCRIPTION: Arg Arq Met Thr Leu Val Phe Gly Ser Pro Arq Asp 5 10

Asp

His 15

- (2) INFORMATION FOR SEQ ID NO: 81:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (1009-1020)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 81:
- Arg Met Thr Leu Val Phe Gly Ser Arg Arg Pro 5
 - (2) INFORMATION FOR SEQ ID NO: 82:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (1009-1017)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 82:
- Arg Met Thr Leu Val Phe Gly Ser Arg-amide 5
 - (2) INFORMATION FOR SEQ ID NO: 83:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (1009-1014)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 83:
- Arg Met Thr Leu Val Phe-amide

- (2) INFORMATION FOR SEQ ID NO: 84:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: TRUNCATED HUMAN iNOS (40-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 84:

Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Lys 5 10

- (2) INFORMATION FOR SEQ ID NO: 85:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: TRUNCATED HUMAN INOS (784-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 85:

Val Gln Gly Ile Leu Glu Arg Val Val 5

- (2) INFORMATION FOR SEQ ID NO: 86:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 86:

Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu
5 10
Ser Lys Gln Gln Asn

Ser Lys Gln Gln Asn 15

(2) INFORMATION FOR SEQ ID NO: 87: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (41-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 87: Gln Asp Asp Leu 5 (2) INFORMATION FOR SEQ ID NO: 88: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (40-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 88: Thr Gln Asp Leu Gln Asp 5 (2) INFORMATION FOR SEQ ID NO: 89: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (39-45)

(B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(D) OTHER INFORMATION: SEQUENCE DESCRIPTION:

Asp

Asp

5

(xi)

Val

Thr

Gln

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

Leu Gln

SEQ ID NO: 89:

- (2) INFORMATION FOR SEQ ID NO: 90:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (38-45)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 90:
- Pro Val Thr Gln Asp Asp Leu Gln 5
 - (2) INFORMATION FOR SEQ ID NO: 91:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-45)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 91:
- Ser Pro Val Thr Gln Asp Asp Leu Gln 5
 - (2) INFORMATION FOR SEQ ID NO: 92:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (40-44)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 92:

Thr Gln Asp Asp Leu

(2) INFORMATION FOR SEQ ID NO: 93: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (39-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 93: Val Thr Gln Asp Asp Leu 5 (2) INFORMATION FOR SEQ ID NO: 94: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (38-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 94: (xi) Val Thr Gln Asp Asp Leu 5 (2) INFORMATION FOR SEQ ID NO: 95: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID-(D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

(xi)

Ser Pro Val

SEQUENCE DESCRIPTION:

Thr

5

Gln Asp Asp Leu

SEQ ID NO: 95:

- (2) INFORMATION FOR SEQ ID NO: 96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (36-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 96: Thr Gln Asp Asp Leu Ser Ser Pro Val 5 (2) INFORMATION FOR SEQ ID NO: 97: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (39-43) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 97: Val Thr Gln Asp Asp (2) INFORMATION FOR SEQ ID NO: 98: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN iNOS (38-43) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- Pro Val Thr Gln Asp Asp

(D) OTHER INFORMATION:

SEQUENCE DESCRIPTION:

o Val Thr Gln Asp

(xi)

SEQ ID NO: 98:

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

- (2) INFORMATION FOR SEQ ID NO: 99:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-43)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

Ser Pro Val Thr Gln Asp Asp 5

- (2) INFORMATION FOR SEQ ID NO: 100:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (36-43)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

Ser Ser Pro Val Thr Gln Asp Asp 5

- (2) INFORMATION FOR SEQ ID NO: 101:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (35-43)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

Thr Ser Ser Pro Val Thr Gln Asp Asp
5

(2) INFORMATION FOR SEQ ID NO: 102: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (37-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: (xi) **SEQ ID NO: 102:** Tyr His Asn Leu Ser Pro Val Thr Gln Asp Asp Leu Gln 5 10 Ser Lys Gln Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 103: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (40-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 103: Leu Gln Tyr His Asn Leu Ser Lys Gln Thr Gln Asp Asp 5 10 Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 104: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (43-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 104:

(xi)

Asp Leu Gln Tyr Gln Gln Asn His Asn Leu Ser Lys 5 10

- (2) INFORMATION FOR SEQ ID NO: 105:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (46-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: **SEQ ID NO: 105:**

Ser Lys Gln Gln Asn Tyr His Asn Leu 5

- (2) INFORMATION FOR SEQ ID NO: 106:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- MOLECULE TYPE: PEPTIDE (ii)
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (49-54)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- SEQUENCE DESCRIPTION: SEQ ID NO: 106: (xi)

Leu Ser Lys Gln Gln 5

- (2) INFORMATION FOR SEQ ID NO: 107:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-51)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:
- Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Pro Val

5 10

Ser Lys

(2) INFORMATION FOR SEQ ID NO: 108:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-48)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:

Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn 5 10

- (2) INFORMATION FOR SEQ ID NO: 109:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-45)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:

Ser Pro Val Thr Gln Asp Asp Leu Gln 5

- (2) INFORMATION FOR SEQ ID NO: 110:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (37-42)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

Ser Pro Val Thr Gln Asp 5

- (2) INFORMATION FOR SEQ ID NO: 111:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (35-44)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

Thr Ser Ser Pro Val Thr Gln Asp Asp Leu
5 10

- (2) INFORMATION FOR SEQ ID NO: 112:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (781-798)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10

- Gly Pro Thr Pro His
 - (2) INFORMATION FOR SEQ ID NO: 113:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (788-792)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113: Val Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 114: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (787-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 114: Val Ile Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 115: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (786-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 115: (xi) Gly Ile Leu Glu Arg Val Val (2) INFORMATION FOR SEQ ID NO: 116: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE

(ix)

FEATURE:

(A) NAME/KEY: HUMAN iNOS (785-792)

(D) OTHER INFORMATION:

(B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:

Gln Gly Ile Leu Glu Arg Val Val

- (2) INFORMATION FOR SEQ ID NO: 117:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (784-792)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117:
- Val Gln Gly Ile Leu Glu Arg Val Val
 - (2) INFORMATION FOR SEQ ID NO: 118:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (787-791)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 118:
- Ile Leu Glu Arg Val
 - (2) INFORMATION FOR SEQ ID NO: 119:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (786-791)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 119: Gly Ile Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 120:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (785-791)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 120:
- Ile Leu Glu Arg Val Gln Gly 5
 - (2) INFORMATION FOR SEQ ID NO: 121:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (784-791)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- SEQUENCE DESCRIPTION: SEQ ID NO: 121: (xi)
- Gln Gly Ile Leu Glu Arg Val Val 5
 - (2) INFORMATION FOR SEQ ID NO: 122:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (783-791)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 122:

Leu Val Gln Gly Ile Leu Glu Arg Val

- (2) INFORMATION FOR SEQ ID NO: 123:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (786-790)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 123:
- Gly Ile Leu Glu Arg 5
 - (2) INFORMATION FOR SEQ ID NO: 124:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (785-790)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 124:
- Gln Gly Ile Leu Glu Arg 5
 - (2) INFORMATION FOR SEQ ID NO: 125:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN INOS (784-790)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 125:

Val Gln Gly Ile Leu Glu Arg 5

- (2) INFORMATION FOR SEQ ID NO: 126:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
 - (A) NAME/KEY: HUMAN iNOS (783-790)
 - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
 - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
 - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 126:

Leu Val Gln Gly Ile Leu Glu Arg